

SEQUENCE LISTING

<110> Roch, Jean-Marc
Bartel, Paul L.
Heichman, Karen

<120> Protein-Protein Interactions in Neurodegenerative
Diseases

<130> Protein Interactions in ND

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<150> US 60/240,790
<151> 2000-10-17

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<170> PatentIn Ver. 2.0

2052660

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<213> Artificial Sequence

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<223> Description of Artificial Sequence:tail for
forward primer for yeast two-hybrid system

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reverse primer for yeast two-hybrid system

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taattaaac atttttcagg gacgttagca tccagagaca ttccatttatt gttccatttga 180
cctttccctc atcaactgagt cctttggagc tgagtt atg tca aca gct gcc tta 234

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Leu Ser Ser Arg Leu Leu Gln Asp Asp Arg Arg Val Thr Pro Thr Cys		
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cac agc tcc act tca gag cct agg tgt tct cgg ttt gac cca gat ggt	378	
His Ser Ser Thr Ser Glu Pro Arg Cys Ser Arg Phe Asp Pro Asp Gly		
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agt ggg agt cca gct acc tgg gac aat ttt ggg atc tgg gat aac cgc	426	
Ser Gly Ser Pro Ala Thr Trp Asp Asn Phe Gly Ile Trp Asp Asn Arg		
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att gat gag cca att ctg ctg cca ccc agc att aag tat ggc aag cca	474	
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Glu Val Leu Tyr Phe Ala Val Tyr Asp Gly His Gly Pro Ala Ala		
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Pro Lys Glu Lys Asn Leu Glu Thr Leu Leu Thr Leu Ala Phe Leu Glu		
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ata gat aaa gcc ttt tcg agt cat gcc cgc ctg tct gct gat gca act	762	
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 Arg Phe Asp Pro Asp Gly Ser Gly Ser Pro Ala Thr Trp Asp Asn Phe
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 65 70 75 80
 Ile Lys Tyr Gly Lys Pro Ile Pro Lys Ile Ser Leu Glu Asn Val Gly
 85 90 95
 Cys Ala Ser Gln Ile Gly Lys Arg Lys Glu Asn Glu Asp Arg Phe Asp
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 Phe Ala Gln Leu Thr Asp Glu Val Leu Tyr Phe Ala Val Tyr Asp Gly
 115 120 125
 His Gly Pro Ala Ala Ala Asp Phe Cys His Thr His Met Glu Lys
 130 135 140
 Cys Ile Met Asp Leu Leu Pro Lys Glu Lys Asn Leu Glu Thr Leu Leu
 145 150 155 160
 Thr Leu Ala Phe Leu Glu Ile Asp Lys Ala Phe Ser Ser His Ala Arg
 165 170 175
 Leu Ser Ala Asp Ala Thr Leu Leu Thr Ser Gly Thr Thr Ala Thr Val
 180 185 190
 Ala Leu Leu Arg Asp Gly Ile Glu Leu Val Val Ala Ser Val Gly Asp
 195 200 205
 Ser Arg Ala Ile Leu Cys Arg Lys Gly Lys Pro Met Lys Leu Thr Ile
 210 215 220

Asp His Thr Pro Glu Arg Lys Asp Glu Lys Glu Arg Ile Lys Lys Cys
 225 230 235 240
 Gly Gly Phe Val Ala Trp Asn Ser Leu Gly Gln Pro His Val Asn Gly
 245 250 255
 Arg Leu Ala Met Thr Arg Ser Ile Gly Asp Leu Asp Leu Lys Thr Ser
 260 265 270
 Gly Val Ile Ala Glu Pro Glu Thr Lys Arg Ile Lys Leu His His Ala
 275 280 285
 Asp Asp Ser Phe Leu Val Leu Thr Thr Asp Gly Ile Asn Phe Met Val
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 Asn Ser Gln Glu Ile Cys Asp Phe Val Asn Gln Cys His Asp Pro Asn
 305 310 315 320
 Glu Ala Ala His Ala Val Thr Glu Gln Ala Ile Gln Tyr Gly Thr Glu
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 Asp Asn Ser Thr Ala Val Val Val Pro Phe Gly Ala Trp Gly Lys Tyr
 340 345 350
 Lys Asn Ser Glu Ile Asn Phe Ser Phe Ser Arg Ser Phe Ala Ser Ser
 355 360 365
 Gly Arg Trp Ala
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